RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _/0/550,584_

Source: _AG/10_

Date Processed by STIC: _/0/5/05_

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Raw Sequence Listing before editing
(for reference only)

PCT

RAW SEQUENCE LISTING   DATE: 10/05/2005
PATENT APPLICATION: US/10/550,584   TIME: 16:38:28

Input Set : A:\Seq Listing as filed 22Sept05.txt
Output Set: N:\CRF4\10052005\J550584.raw

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  3 <110> APPLICANT: SHINOZAKI, Kazuko
  4         KATSURA, Koji
  5         ITO, Yusuke
  7 <120> TITLE OF INVENTION: Stress-inducible promoter and method for using the same
  9 <130> FILE REFERENCE: 382.1048
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/550,584
C--> 12 <141> CURRENT FILING DATE: 2005-09-22
 14 <150> PRIOR APPLICATION NUMBER: JP 2003-080847
 15 <151> PRIOR FILING DATE: 2003-03-24
 17 <160> NUMBER OF SEQ ID NOS: 12
 19 <170> SOFTWARE: PatentIn Ver. 2.1
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ERRORED SEQUENCES

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216 <210> SEQ ID NO: 7
217 <211> LENGTH: 1437
218 <212> TYPE: DNA
219 <213> ORGANISM: Arabidopsis thaliana
221 <220> FEATURE:
222 <221> NAME/KEY: CDS
223 <222> LOCATION: (167)..(1171)
225 <400> SEQUENCE: 7
226 gctgtctgat aaaaagaaga ggaaaactcg aaaaagctac acacaagaag aagaagaaaa 60
227 gatacgagca agaagactaa acacgaaagc gatttatcaa ctcgaaggaa gagactttga 120
228 ttttcaaatt tcgtccccta tagattgtgt tgtttctggg aaggag atg gca gtt   175
229                                                        Met Ala Val
230                                                        1
231 tat gat cag agt gga gat aga aac aga aca caa att gat aca tcg agg   223
232 Tyr Asp Gln Ser Gly Asp Arg Asn Arg Thr Gln Ile Asp Thr Ser Arg
233      5                  10                  15
234 aaa agg aaa tct aga agt aga ggt gac ggt act act gtg gct gag aga   271
235 Lys Arg Lys Ser Arg Ser Arg Gly Asp Gly Thr Thr Val Ala Glu Arg
236  20                  25                  30                  35
237 tta aag aga tgg aaa gag tat aac gag acc gta gaa gaa gtt tct acc   319
238 Leu Lys Arg Trp Lys Glu Tyr Asn Glu Thr Val Glu Glu Val Ser Thr
239                  40                  45                  50
240 aag aag agg aaa gta cct gcg aaa ggg tcg aag aag ggt tgt atg aaa   367
241 Lys Lys Arg Lys Val Pro Ala Lys Gly Ser Lys Lys Gly Cys Met Lys
242                  55                  60                  65
243 ggt aaa gga gga cca gag aat agc cga tgt agt ttc aga gga gtt agg   415
244 Gly Lys Gly Gly Pro Glu Asn Ser Arg Cys Ser Phe Arg Gly Val Arg
245                  70                  75                  80
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246 caa agg att tgg ggt aaa tgg gtt gct gag atc aga gag cct aat cga   463
247 Gln Arg Ile Trp Gly Lys Trp Val Ala Glu Ile Arg Glu Pro Asn Arg
248        85              90                      95
249 ggt agc agg ctt tgg ctt ggt act ttc cct act gct caa gaa gct gct   511
250 Gly Ser Arg Leu Trp Leu Gly Thr Phe Pro Thr Ala Gln Glu Ala Ala
251 100              105                     110                   115
252 tct gct tat gat gag gct gct aaa gct atg tat ggt cct ttg gct cgt   559
253 Ser Ala Tyr Asp Glu Ala Ala Lys Ala Met Tyr Gly Pro Leu Ala Arg
254              120                     125                   130
255 ctt aat ttc cct cgg tct gat gcg tct gag gtt acg agt acc tca agt   607
256 Leu Asn Phe Pro Arg Ser Asp Ala Ser Glu Val Thr Ser Thr Ser Ser
257              135                     140                   145
258 cag tct gag gtg tgt act gtt gag act cct ggt tgt gtt cat gtg aaa   655
259 Gln Ser Glu Val Cys Thr Val Glu Thr Pro Gly Cys Val His Val Lys
260              150                     155                   160
261 aca gag gat cca gat tgt gaa tct aaa ccc ttc tcc ggt gga gtg gag   703
262 Thr Glu Asp Pro Asp Cys Glu Ser Lys Pro Phe Ser Gly Gly Val Glu
263              165                     170                   175
264 ccg atg tat tgt ctg gag aat ggt gcg gaa gag atg aag aga ggt gtt   751
265 Pro Met Tyr Cys Leu Glu Asn Gly Ala Glu Glu Met Lys Arg Gly Val
266 180              185                     190                   195
267 aaa gcg gat aag cat tgg ctg agc gag ttt gaa cat aac tat tgg agt   799
268 Lys Ala Asp Lys His Trp Leu Ser Glu Phe Glu His Asn Tyr Trp Ser
269              200                     205                   210
270 gat att ctg aaa gag aaa gag aaa cag aag gag caa ggg att gta gaa   847
271 Asp Ile Leu Lys Glu Lys Glu Lys Gln Lys Glu Gln Gly Ile Val Glu
272              215                     220                   225
273 acc tgt cag caa caa cag cag gat tcg cta tct gtt gca gac tat ggt   895
274 Thr Cys Gln Gln Gln Gln Gln Asp Ser Leu Ser Val Ala Asp Tyr Gly
275              230                     235                   240
276 tgg ccc aat gat gtg gat cag agt cac ttg gat tct tca gac atg ttt   943
277 Trp Pro Asn Asp Val Asp Gln Ser His Leu Asp Ser Ser Asp Met Phe
278        245                     250                     255
279 gat gtc gat gag ctt cta cgt gac cta aat ggc gac gat gtg ttt gca   991
280 Asp Val Asp Glu Leu Leu Arg Asp Leu Asn Gly Asp Asp Val Phe Ala
281 260                     265                     270           275
282 ggc tta aat cag gac cgg tac ccg ggg aac agt gtt gcc aac ggt tca   1039
283 Gly Leu Asn Gln Asp Arg Tyr Pro Gly Asn Ser Val Ala Asn Gly Ser
284              280                     285                   290
285 tac agg ccc gag agt caa caa agt ggt ttt gat ccg cta caa agc ctc   1087
286 Tyr Arg Pro Glu Ser Gln Gln Ser Gly Phe Asp Pro Leu Gln Ser Leu
287              295                     300                   305
288 aac tac gga ata cct ccg ttt cag ctc gag gga aag gat ggt aat gga   1135
289 Asn Tyr Gly Ile Pro Pro Phe Gln Leu Glu Gly Lys Asp Gly Asn Gly
290              310                     315                   320
291 ttc ttc gac gac ttg agt tac ttg gat ctg gag aac taaacaaac   1181
292 Phe Phe Asp Asp Leu Ser Tyr Leu Asp Leu Glu Asn
293 325                     330                     335
294 aatatgaagc tttttggatt tgatatttgc cttaatccca caacgactgt tgattctcta   1241
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    295 tccgagtttt agtgatatag agaactacag aacacgtttt ttcttgttat aaaggtgaac 1301
    296 tgtatatatc gaaacagtga tatgacaata gagaagacaa ctatagtttg ttagtctgct 1361
    297 tctcttaagt tgttctttag atatgtttta tgttttgtaa caacaggaat gaataataca 1421
E--> 298 cacttgtaaa aaaaaa   1437 ← insert

VERIFICATION SUMMARY   DATE: 10/05/2005
PATENT APPLICATION: US/10/550,584   TIME: 16:38:29

Input Set : A:\Seq Listing as filed 22Sept05.txt
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L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:148 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=6
L:298 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:1437 SEQ:7